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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CC genome; cellular implants comprising a TC transfected with (I); solated (I); and transgenic animals expressing (A). Activation of cellular product, particularly a heterologous therapeutic protein encoded by a second inserted nucleic acid sequence. Particularly it is used to expand the relatively few cells that are successfully transfected during gene therapy procedures. Other responses that can be regulated are cell migration and contraction, or pigment production. In transgenic animals, expression or stimulation of (A) is designed to develop cardiac arrhythmia, symptoms of bone disease, setzures, vascular contractions, dementia, neurodegeneration etc., for use as models content in eggshells or altered fat/lean ratios) or to control for production of improved food products (e.g. increased calcium content in eggshells or altered fat/lean ratios) or to control fertility or induce labour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 252;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                           15-SEP-1994 (first entry)
Partial sequence of the human kappa opioid receptor genomic clone H14 (KORA).
                                                                                                                                                                                                                                                                                                                                                                                                                                                   056702 standard; DNA; 2447 BP. Sequ-Incl Comports on A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               human KOR cDNA (see T90998) and includes flanking sequences that encode signal and epitope peptides to facilitate the detection and purification of recombinant OR1. A novel method for selectively activating a target cell (TC) comprises: (1) introducing into the cell a nucleic acid sequence (I) that expresses a RASSI (A) and (1i) exposing the transfected cell to small synthetic molecules (B) that bind to and activate (A), inducing the G protein coupled cellular response associated with receptor activation. (A) has: (a) decreased binding affinity for a selected natural ligand of the native receptor; (b) binding affinity for [B); and (c) is activated by binding (B) sufficiently to produce the required cellular response. Also new are: transgenic cells including heterologous (I) in the
WPI; 94\text{-}08399\text{/}10. DNA encoding opioid receptors and antibodies against this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1143 ACTGAAGATGAG 1154
                                                                                                 Edwards RH,
                                                                                                                              (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1083 CATTCTCTACGCCTTTCTTGATGAAAACTTCAAGCGGTGTTTCCGGGGACTTCTGCTTTCC 1142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1023 TGCTCTCCAGCTATTACTTCTGCATCGCCTTAGGCTATACCAACAGTAGCCTGAATCC 1082
                                                                                                                                                                                                                                  03-MAR-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sequence
                                                                                                                                                                                                                                                                                                                            Opioid receptor; morphine; opiate; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        884 ACTGAAGATGAG 895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  824 CATTCTCTACGCCTTTCTTGATGAAAACTTCAAGCGGTGTTTCCCGGGACTTCTGCTTTCC 883
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          903 CAACCIGCGTAGGATCACCAGACIGGTCCTGGTGGTGGTGGCAGTCTTCGTCGTCTGCTG 962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  644 CAACCIGCGTAGGATCACCAGACIGGTCCTGGTGGTGGTGGCAGTCTTCGTCGTCTGCTG 703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            963 GACTCCCATTCACATATTCATCCTGGTGGAGGCTCTGGGGAGCACCTCCCACAGCAGC 1022
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      704 GACTCCCATTCACATATTCATCCTGGTGGAGGCTCTGGGGAGCACCTCCCACAGCACAGC 763
                                                                                                                                                                                              AUG-1993; U07665
                                                                94-083099/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1275 BP;
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                                                                                                 Evans CJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        25.2%; Score 252; DB 1; L
100.0%; Pred. No. 2.4e-116;
tive 0; Mismatches 0;
                                                                                                 Kaufman D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          266 A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          373 C;
                                                                                                    Keith DE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          318 G;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 1275;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          318 T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Activation of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              В
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The nucleotide sequence of the novel mouse kappa opioid receptor gene mORKI. The gene was isolated from a mouse brain cDNA library using a fragment (amplified from the cDNA library with primers 075929-30) as a probe. The primers are based on the conserved sequences present in the second and third transmembrane domains of somatostatin (SRIF) receptor subtypes SSTR1, SSTR2 and SSTR3. The 1.2 kb PstI fragment from the mous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Polynucleotides and peptides derived from opioid recept polypeptides - fc use in therapewtic compositions and
                                                                                                                                                                                                                                                                                                                                                                Bell GI, Reisine T, Yasuda K; WPI; 95-022804/03.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-JUL-1993;
05-NOV-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mouse; kappa; delta; mu; opioid receptor; brain; primer; PCR; amplify; transmembrane domain; somatostatin; receptor; human; expression vector;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-AUG-1995 (first entry)
Mouse kappa opioid receptor mork1 cDNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     receptor - used to express and locate these receptors, and screen cpds. for opioid (anniagonist activity Example; Fig 8b; 74pp; English.
To isolate opiate receptor genomic clones, 300,000 human genomic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            08-DEC-1994.
20-MAY-1994; U05747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO9428132-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      truncate; chimaeric; assay; probe; ss
                                                                                                                                                                                                                                                                                                                                                                                                                         (ARCH-) ARCH DEV CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            275926 standard; DNA; 1410 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          451 AAGGT 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181 GGCTGCTGTCGTCATCTGTTGGCATCTCTGCAATAGTCCTTGGAGGCACCAAAGTCAGGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          391 GGCTGCTGTCGTCATCTGTTGGCATCTC° CAATAGTCCTTGGAGGCACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         331 CCGTGAAGGCTTT:GACTTCCGCACACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          211 CCTGGCCTTTTGGGGATGTGCTGTGCAAGATAGTAATTTCCATTGATTACTACAACATGT 270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     121 CCG1GAAGGCT11GACTTCCGCACACC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       271 TCACCAGCATCTTCACCTTGACCATGATGAGCGTGGACCGCTACATTGCCC...gTGCCACC 330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 TCACCAGCATCTTCACCTTGACCATGATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 CCTGGCCTTTTGGGGATGTGCTGCAAGATAGTAATTTCCATTGATTACTACAACATGT 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-066296.
US-100694.
US-147592.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /product= mouse kappa opioid receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 245; DB 1; Length . . ?; Pred No. 7.5e-113;
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